

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/791,373

Source: _____

Date Processed by STIC: _____

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IFWO

RAW SEQUENCE LISTING

DATE: 04/20/2005

PATENT APPLICATION: US/10/791,373

TIME: 10:55:43

Input Set : N:\Cr3\RULE60\10791373.raw.txt

Output Set: N:\CRF4\04202005\J791373.raw

SEQUENCE LISTING

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3 (1) GENERAL INFORMATION:
5     (i) APPLICANT: Chee, Mark S.
6             Wang, Chunwei
7             Jevons, Luis C.
8             Bernhart, Derek H.
9             Lipshutz, Robert J.
11    (ii) TITLE OF INVENTION: Computer-Aided Visualization and
12                             Analysis System for Sequence Evaluation
14    (iii) NUMBER OF SEQUENCES: 39
16    (iv) CORRESPONDENCE ADDRESS:
17            (A) ADDRESSEE: Townsend and Townsend and Crew LLP
18            (B) STREET: Two Embarcadero Center, 8th Floor
19            (C) CITY: San Francisco
20            (D) STATE: California
21            (E) COUNTRY: USA
22            (F) ZIP: 94111-3834
24    (v) COMPUTER READABLE FORM:
25            (A) MEDIUM TYPE: Floppy disk
26            (B) COMPUTER: IBM PC compatible
27            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
28            (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
30    (vi) CURRENT APPLICATION DATA:
C--> 31            (A) APPLICATION NUMBER: US/10/791,373
C--> 32            (B) FILING DATE: 01-Mar-2004
W--> 33            (C) CLASSIFICATION: 435
35    (vii) PRIOR APPLICATION DATA:
36            (A) APPLICATION NUMBER: US/08/327,525
37            (B) FILING DATE: October 21, 1994
40    (viii) ATTORNEY/AGENT INFORMATION:
41            (A) NAME: Norviel, Vernon A.
42            (B) REGISTRATION NUMBER: 32,483
43            (C) REFERENCE/DOCKET NUMBER: 16528X-82
45    (ix) TELECOMMUNICATION INFORMATION:
46            (A) TELEPHONE: 415-326-2400
47            (B) TELEFAX: 415-326-2422
50 (2) INFORMATION FOR SEQ ID NO: 1:
52     (i) SEQUENCE CHARACTERISTICS:
53            (A) LENGTH: 15 base pairs
54            (B) TYPE: nucleic acid
55            (C) STRANDEDNESS: single
56            (D) TOPOLOGY: linear
W--> 58     (ii) MOLECULE TYPE: DNA (oligonucleotide)

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62      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
64 ATGTGGACAG TTGTA                                     15
66 (2) INFORMATION FOR SEQ ID NO: 2:
68      (i) SEQUENCE CHARACTERISTICS:
69          (A) LENGTH: 15 base pairs
70          (B) TYPE: nucleic acid
71          (C) STRANDEDNESS: single
72          (D) TOPOLOGY: linear
W--> 74      (ii) MOLECULE TYPE: DNA (oligonucleotide)
78      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
80 ATGTGGATAG TTGTA                                     15
82 (2) INFORMATION FOR SEQ ID NO: 3:
84      (i) SEQUENCE CHARACTERISTICS:
85          (A) LENGTH: 15 base pairs
86          (B) TYPE: nucleic acid
87          (C) STRANDEDNESS: single
88          (D) TOPOLOGY: linear
W--> 90      (ii) MOLECULE TYPE: DNA (oligonucleotide)
94      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
96 ATGTGGAKAG TTGTA                                     15
98 (2) INFORMATION FOR SEQ ID NO: 4:
100      (i) SEQUENCE CHARACTERISTICS:
101          (A) LENGTH: 11 base pairs
102          (B) TYPE: nucleic acid
103          (C) STRANDEDNESS: single
104          (D) TOPOLOGY: linear
W--> 106      (ii) MOLECULE TYPE: DNA (oligonucleotide)
110      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
112 AAAACTGAAA A                                         11
114 (2) INFORMATION FOR SEQ ID NO: 5:
116      (i) SEQUENCE CHARACTERISTICS:
117          (A) LENGTH: 11 base pairs
118          (B) TYPE: nucleic acid
119          (C) STRANDEDNESS: single
120          (D) TOPOLOGY: linear
W--> 122      (ii) MOLECULE TYPE: DNA (oligonucleotide)
126      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
128 AAAACCGAAA A                                         11
130 (2) INFORMATION FOR SEQ ID NO: 6:
132      (i) SEQUENCE CHARACTERISTICS:
133          (A) LENGTH: 17 base pairs
134          (B) TYPE: nucleic acid
135          (C) STRANDEDNESS: single
136          (D) TOPOLOGY: linear
W--> 138      (ii) MOLECULE TYPE: DNA (oligonucleotide)
142      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
144 AAACCCAATC CACATCA                                   17
146 (2) INFORMATION FOR SEQ ID NO: 7:
148      (i) SEQUENCE CHARACTERISTICS:

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149          (A) LENGTH: 17 base pairs
150          (B) TYPE: nucleic acid
151          (C) STRANDEDNESS: single
152          (D) TOPOLOGY: linear
W--> 154      (ii) MOLECULE TYPE: DNA (oligonucleotide)
158          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
160 AAACCCAGTC CACATCA                                     17
162 (2) INFORMATION FOR SEQ ID NO: 8:
164      (i) SEQUENCE CHARACTERISTICS:
165          (A) LENGTH: 31 base pairs
166          (B) TYPE: nucleic acid
167          (C) STRANDEDNESS: single
168          (D) TOPOLOGY: linear
W--> 170      (ii) MOLECULE TYPE: DNA (oligonucleotide)
174          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
176 GGGGAAGCAG ATTTGGGTAC CACCCAAGTA T                     31
178 (2) INFORMATION FOR SEQ ID NO: 9:
180      (i) SEQUENCE CHARACTERISTICS:
181          (A) LENGTH: 31 base pairs
182          (B) TYPE: nucleic acid
183          (C) STRANDEDNESS: single
184          (D) TOPOLOGY: linear
W--> 186      (ii) MOLECULE TYPE: DNA (oligonucleotide)
190          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
192 GGGGAAGCAG ATTTGAAMAC CACCCAAGTA T                     31
194 (2) INFORMATION FOR SEQ ID NO: 10:
196      (i) SEQUENCE CHARACTERISTICS:
197          (A) LENGTH: 59 base pairs
198          (B) TYPE: nucleic acid
199          (C) STRANDEDNESS: single
200          (D) TOPOLOGY: linear
W--> 202      (ii) MOLECULE TYPE: DNA (oligonucleotide)
206          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
208 GCATTAGTAG AGATATGTAC AGAAATGGAA AAGGAAGGGA AAATTTCAAA AATTGGGCC   59
210 (2) INFORMATION FOR SEQ ID NO: 11:
212      (i) SEQUENCE CHARACTERISTICS:
213          (A) LENGTH: 59 base pairs
214          (B) TYPE: nucleic acid
215          (C) STRANDEDNESS: single
216          (D) TOPOLOGY: linear
W--> 218      (ii) MOLECULE TYPE: DNA (oligonucleotide)
222          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
224 GCATTAGTAG AAATTTGTAC AGAGATGGAA AAGGAAGGGA AAATTTCAAA AATTGGGCC   59
226 (2) INFORMATION FOR SEQ ID NO: 12:
228      (i) SEQUENCE CHARACTERISTICS:
229          (A) LENGTH: 59 base pairs
230          (B) TYPE: nucleic acid
231          (C) STRANDEDNESS: single
232          (D) TOPOLOGY: linear

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W--> 234      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      238      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
      240 GCATTAGTAG AGATATGGAG AGRARDGGRA ANNNAAGGGA AAATTNNNAA AATTGGGCC      59
      242 (2) INFORMATION FOR SEQ ID NO: 13:
      244      (i) SEQUENCE CHARACTERISTICS:
      245          (A) LENGTH: 59 base pairs
      246          (B) TYPE: nucleic acid
      247          (C) STRANDEDNESS: single
      248          (D) TOPOLOGY: linear

W--> 250      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      254      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
      256 GCATTAGTAG AGATATGKAS AGRARDGGRA ANNNAAGGGA AAATNNNAA AATTGGGCC      59
      259 (2) INFORMATION FOR SEQ ID NO: 14:
      261      (i) SEQUENCE CHARACTERISTICS:
      262          (A) LENGTH: 59 base pairs
      263          (B) TYPE: nucleic acid
      264          (C) STRANDEDNESS: single
      265          (D) TOPOLOGY: linear

W--> 267      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      271      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
      273 GCATTAGTAG AGATATGKAS AGRRRDGGRA ANNNAAGGGA AAADTYNNAA AATTGGGCC      59
      275 (2) INFORMATION FOR SEQ ID NO: 15:
      277      (i) SEQUENCE CHARACTERISTICS:
      278          (A) LENGTH: 59 base pairs
      279          (B) TYPE: nucleic acid
      280          (C) STRANDEDNESS: single
      281          (D) TOPOLOGY: linear

W--> 283      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      287      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
      289 GCATTAGTAG AGATATGTAS AGRRADGGAA ANGGAAGGGA AAATTNNNNA AATTGGGCC      59
      291 (2) INFORMATION FOR SEQ ID NO: 16:
      293      (i) SEQUENCE CHARACTERISTICS:
      294          (A) LENGTH: 59 base pairs
      295          (B) TYPE: nucleic acid
      296          (C) STRANDEDNESS: single
      297          (D) TOPOLOGY: linear

W--> 299      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      303      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
      305 GCATTAGTAG AGATATGTAC AGRGAGGGAA ANGGAAGGGA AAATTNNNNA AATTGGGCC      59
      307 (2) INFORMATION FOR SEQ ID NO: 17:
      309      (i) SEQUENCE CHARACTERISTICS:
      310          (A) LENGTH: 59 base pairs
      311          (B) TYPE: nucleic acid
      312          (C) STRANDEDNESS: single
      313          (D) TOPOLOGY: linear

W--> 315      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      319      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
      321 GCATTAGTAG AGATATGTAS AGRGAGGGAA ANGGAAGGGA AAATTNNNNA AATTGGGCC      59
      323 (2) INFORMATION FOR SEQ ID NO: 18:

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Output Set: N:\CRF4\04202005\J791373.raw

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325      (i) SEQUENCE CHARACTERISTICS:
326          (A) LENGTH: 59 base pairs
327          (B) TYPE: nucleic acid
328          (C) STRANDEDNESS: single
329          (D) TOPOLOGY: linear
W--> 331      (ii) MOLECULE TYPE: DNA (oligonucleotide)
335          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
337 GCATTAGTAG GAGGNNNGAC AGGGRKGGAA ANNMAAGGGA AAATNNNAA AATTGGGCC      59
339 (2) INFORMATION FOR SEQ ID NO: 19:
341      (i) SEQUENCE CHARACTERISTICS:
342          (A) LENGTH: 160 base pairs
343          (B) TYPE: nucleic acid
344          (C) STRANDEDNESS: single
345          (D) TOPOLOGY: linear
W--> 347      (ii) MOLECULE TYPE: DNA (oligonucleotide)
351          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
353 TCGAGATAAT CTATGTCCTC GTCTACTATG TCATAATCTT CTTTACTTAA ACGGTCCTTT      60
355 TACCTTTGGT TTTTACTATC CCCCTTAACC TCCAAAATAG TTTCATTCTG TCATGCTAGT      120
357 CTATGGACAT CTTTAGACAC CTGTATTTTCG ATATCCATGT      160
359 (2) INFORMATION FOR SEQ ID NO: 20:
361      (i) SEQUENCE CHARACTERISTICS:
362          (A) LENGTH: 160 base pairs
363          (B) TYPE: nucleic acid
364          (C) STRANDEDNESS: single
365          (D) TOPOLOGY: linear
W--> 367      (ii) MOLECULE TYPE: DNA (oligonucleotide)
371          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
373 NNGAGATANN NTATGTCCTC GTCYACTATG TNANNNNNNNN NNNNNNNNAA ACGGTCCTNN      60
375 NNNNNNNNNN NNNNNNNNNN CNNCNTAACC TCCAAAATAN NNNNNNTCTN NNNNANNNT      120
377 CTANNNGNAG NNNNAGANAR NCCNNNNNNN NNATNCATGT      160
379 (2) INFORMATION FOR SEQ ID NO: 21:
381      (i) SEQUENCE CHARACTERISTICS:
382          (A) LENGTH: 160 base pairs
383          (B) TYPE: nucleic acid
384          (C) STRANDEDNESS: single
385          (D) TOPOLOGY: linear
W--> 387      (ii) MOLECULE TYPE: DNA (oligonucleotide)
391          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
393 TCGAGATAAT CTATGTCCTC GTCTACTATG TCATAATNNN NNNNACTTAA ACGGTCCTTT      60
395 TACCTTTGGT TTTTACTATC CCCCTTAACC TCCAAAATAG TTTCATTCTG NCATANNAGT      120
397 CTATGNGNNG NNNTAGACAG NCCNNNNNTCG ATATCCATGT      160
399 (2) INFORMATION FOR SEQ ID NO: 22:
401      (i) SEQUENCE CHARACTERISTICS:
402          (A) LENGTH: 160 base pairs
403          (B) TYPE: nucleic acid
404          (C) STRANDEDNESS: single
405          (D) TOPOLOGY: linear
W--> 407      (ii) MOLECULE TYPE: DNA (oligonucleotide)
411          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

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VERIFICATION SUMMARY

DATE: 04/20/2005

PATENT APPLICATION: US/10/791,373

TIME: 10:55:44

Input Set : N:\CrF3\RULE60\10791373.raw.txt

Output Set: N:\CRF4\04202005\J791373.raw

L:31 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:32 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:38 M:238 W: Alpha Fields not Ordered, Reordered [(C) CLASSIFICATION:] of (1) (vi)
L:58 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:74 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:90 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:106 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:122 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5
L:138 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6
L:154 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7
L:170 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8
L:186 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9
L:202 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10
L:218 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11
L:234 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12
L:250 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:267 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14
L:283 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15
L:299 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16
L:315 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17
L:331 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18
L:347 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=19
L:367 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20
L:387 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21
L:407 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22
L:427 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23
L:447 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24
L:467 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:487 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:507 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:527 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:543 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:559 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:575 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:591 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:607 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:623 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34
L:639 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:655 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:671 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:687 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:703 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39